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RAW SEQUENCE LISTING

DATE: 04/01/2003

PATENT APPLICATION: US/10/089,825

TIME: 12:45:51

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Output Set: N:\CRF4\04012003\J089825.raw

1 <110> APPLICANT: Teem, John L.

2 <120> TITLE OF INVENTION: Materials and Methods for Detecting Interaction of CFTR

Polypeptides

3 <130> FILE REFERENCE: FSU-100C2XC1

C--> 4 <140> CURRENT APPLICATION NUMBER: US/10/089,825

5 <141> CURRENT FILING DATE: 2003-01-10

6 <150> PRIOR APPLICATION NUMBER: 60/157,996

7 <151> PRIOR FILING DATE: 1999-10-06

8 <150> PRIOR APPLICATION NUMBER: 60/181,892

9 <151> PRIOR FILING DATE: 2000-02-11

10 <150> PRIOR APPLICATION NUMBER: 60/182,373

11 <151> PRIOR FILING DATE: 2000-02-14

12 <160> NUMBER OF SEQ ID NOS: 4

13 <170> SOFTWARE: PatentIn version 3.0

15 <210> SEQ ID NO: 1

16 <211> LENGTH: 25

17 <212> TYPE: DNA

18 <213> ORGANISM: Homo sapien

19 <400> SEQUENCE: 1

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23 <211> LENGTH: 28

24 <212> TYPE: DNA

25 <213> ORGANISM: Homo sapien

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29 <210> SEQ ID NO: 3

30 <211> LENGTH: 4443

31 <212> TYPE: DNA

32 <213> ORGANISM: Homo sapien

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34 <221> NAME/KEY: gene

35 <222> LOCATION: (1)..(4443)

36 <400> SEQUENCE: 3

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39 ccttctgttg attctgctga caatctatct gaaaaatttg aaagagaatg ggatagagag 180
40 ctggcttcaa agaaaaatcc taaactcatt aatgcccttc ggcatgttt tttctggaga 240
41 tttatgttct atggaatctt tttatattta ggggaagtca ccaaagcagt acagcctctc 300
42 ttactgggaa gaatcatagc ttcctatgac ccggaataca aggaggaacg ctctatcgcg 360
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113 <210> SEQ ID NO: 4

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115 <212> TYPE: PRT

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117 <220> FEATURE:

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120 <400> SEQUENCE: 4

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124 20 25 30
125 Glu Leu Ser Asp Ile Tyr Gln Ile Pro Ser Val Asp Ser Ala Asp Asn
126 35 40 45
127 Leu Ser Glu Lys Leu Glu Arg Glu Trp Asp Arg Glu Leu Ala Ser Lys
128 50 55 60
129 Lys Asn Pro Lys Leu Ile Asn Ala Leu Arg Arg Cys Phe Phe Trp Arg
130 65 70 75 80
131 Phe Met Phe Tyr Gly Ile Phe Leu Tyr Leu Gly Glu Val Thr Lys Ala
132 85 90 95
133 Val Gln Pro Leu Leu Leu Gly Arg Ile Ile Ala Ser Tyr Asp Pro Asp
134 100 105 110
135 Asn Lys Glu Glu Arg Ser Ile Ala Ile Tyr Leu Gly Ile Gly Leu Cys
136 115 120 125
137 Leu Leu Phe Ile Val Arg Thr Leu Leu Leu His Pro Ala Ile Phe Gly
138 130 135 140
139 Leu His His Ile Gly Met Gln Met Arg Ile Ala Met Phe Ser Leu Ile
140 145 150 155 160
141 Tyr Lys Lys Thr Leu Lys Leu Ser Ser Arg Val Leu Asp Lys Ile Ser
142 165 170 175
143 Ile Gly Gln Leu Val Ser Leu Leu Ser Asn Asn Leu Asn Lys Phe Asp
144 180 185 190
145 Glu Gly Leu Ala Leu Ala His Phe Val Trp Ile Ala Pro Leu Gln Val

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149	Cys	Gly	Leu	Gly	Phe	Leu	Ile	Val	Leu	Ala	Leu	Phe	Gln	Ala	Gly	Leu			
150		225				230						235				240			
151	Gly	Arg	Met	Met	Met	Lys	Tyr	Arg	Asp	Gln	Arg	Ala	Gly	Lys	Ile	Ser			
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154			260						265					270					
155	Lys	Ala	Tyr	Cys	Trp	Glu	Glu	Ala	Met	Glu	Lys	Met	Ile	Glu	Asn	Leu			
156		275						280					285						
157	Arg	Gln	Thr	Glu	Leu	Lys	Leu	Thr	Arg	Lys	Ala	Ala	Tyr	Val	Arg	Tyr			
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159	Phe	Asn	Ser	Ser	Ala	Phe	Phe	Phe	Ser	Gly	Phe	Phe	Val	Val	Phe	Leu			
160		305				310					315					320			
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165	Gln	Phe	Pro	Trp	Ala	Val	Gln	Thr	Trp	Tyr	Asp	Ser	Leu	Gly	Ala	Ile			
166		355						360				365							
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172				405						410					415				
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182				485						490					495				
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184			500						505				510						
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186		515						520					525						
187	Asp	Ile	Ser	Lys	Phe	Ala	Glu	Lys	Asp	Asn	Ile	Val	Leu	Gly	Glu	Gly			
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189	Gly	Ile	Thr	Leu	Ser	Gly	Gln	Arg	Ala	Arg	Ile	Ser	Leu	Ala	Arg				
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200              625              630              635              640
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206              675              680              685
207  Gln Thr Gly Glu Phe Gly Glu Lys Arg Lys Asn Ser Ile Leu Asn Pro
208              690              695              700
209  Ile Asn Ser Ile Arg Lys Phe Ser Ile Val Gln Lys Thr Pro Leu Gln
210              705              710              715              720
211  Met Asn Gly Ile Glu Glu Asp Ser Asp Glu Pro Leu Glu Arg Arg Leu
212              725              730              735
213  Ser Leu Val Pro Asp Ser Glu Gln Gly Glu Ala Ile Leu Pro Arg Ile
214              740              745              750
215  Ser Val Ile Ser Thr Gly Pro Thr Leu Gln Ala Arg Arg Arg Gln Ser
216              755              760              765
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219  Arg Lys Thr Thr Ala Ser Thr Arg Lys Val Ser Leu Ala Pro Gln Ala
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222              805              810              815
223  Gly Leu Glu Ile Ser Glu Glu Ile Asn Glu Glu Asp Leu Lys Glu Cys
224              820              825              830
225  Leu Phe Asp Asp Met Glu Ser Ile Pro Ala Val Thr Thr Trp Asn Thr
226              835              840              845
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228              850              855              860
229  Trp Cys Leu Val Ile Phe Leu Ala Glu Val Ala Ala Ser Leu Val Val
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236              915              920              925
237  Met Gly Phe Phe Arg Gly Leu Pro Leu Val His Thr Leu Ile Thr Val
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VERIFICATION SUMMARY

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